

```

QY 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
DB 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
DB 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
QY 241 ASIYQMLDRRSORSERCEVEIP 263
DB 241 ASIYQMLDRRSORSERCEVEIP 263

```

RESULT 2

```

US-10-293-832-28
; Sequence 28, Application US/10293832
; Publication No. US20030180752A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Wei
; APPLICANT: Fouser, Lynette
; APPLICANT: Spaulding, Vikki
; TITLE OF INVENTION: TYPE 2 CYTOKINE RECEPTOR AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 22058-546
; CURRENT APPLICATION NUMBER: US/10/293,832
; PRIOR FILING DATE: 2002-11-12
; PRIOR APPLICATION NUMBER: US 60/332,366
; PRIOR FILING DATE: 2001-11-09
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: Patent in version 3.2
; SEQ ID NO 28
; LENGTH: 263
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-293-832-28

```

```

Query Match 100.0%; Score 1432; DB 12; Length 263;
Best Local Similarity 100.0%; Pred. No. 4e-138;
Matches 263; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 MPPKHCFGLFLISFFLTGAGTQSTHESLKQVQFQSENFNHILQWQGRALTNSSVY 60
DB 1 MPPKHCFGLFLISFFLTGAGTQSTHESLKQVQFQSENFNHILQWQGRALTNSSVY 60
QY 61 FVOYKIMFSCMSKSHQKPSGCHISCNPPGCRITLAKYQKQWKNKDCWGTQELSCDL 120
DB 61 FVOYKIMFSCMSKSHQKPSGCHISCNPPGCRITLAKYQKQWKNKDCWGTQELSCDL 120
QY 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
DB 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
DB 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
QY 241 ASIYQMLDRRSORSERCEVEIP 263
DB 241 ASIYQMLDRRSORSERCEVEIP 263

```

RESULT 3

```

US-10-312-088-42
; Sequence 42, Application US/10312088
; Publication No. US20030219862A1
; GENERAL INFORMATION:
; APPLICANT: Agarwal, Pankaj
; APPLICANT: Cogswell, John P.
; APPLICANT: Kabnic, Karen S.
; APPLICANT: Lai, Ying-Ta
; APPLICANT: Martensen, Shelby A.
; APPLICANT: Murdock, Paul R.
; APPLICANT: Smith, Randall F.

```

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; APPLICANT: Strum, Jay C.
; APPLICANT: Xiang, Zhaoying
; APPLICANT: Xie, Qing
; APPLICANT: Rizni, Safia K.
; TITLE OF INVENTION: NOVEL COMPOUNDS
; FILE REFERENCE: GP50029
; CURRENT APPLICATION NUMBER: US/10/312,088
; CURRENT FILING DATE: 2002-12-20
; PRIOR APPLICATION NUMBER: PCT/US01/19929
; PRIOR FILING DATE: 2001-06-22
; PRIOR APPLICATION NUMBER: 60/213,161
; PRIOR FILING DATE: 2000-06-22
; PRIOR APPLICATION NUMBER: 60/213,156
; PRIOR FILING DATE: 2000-06-22
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: Fast-SEQ for Windows Version 4.0
; SEQ ID NO 42
; LENGTH: 263
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-312-088-42

```

```

Query Match 100.0%; Score 1432; DB 12; Length 263;
Best Local Similarity 100.0%; Pred. No. 4e-138;
Matches 263; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 MPPKHCFGLFLISFFLTGAGTQSTHESLKQVQFQSENFNHILQWQGRALTNSSVY 60
DB 1 MPPKHCFGLFLISFFLTGAGTQSTHESLKQVQFQSENFNHILQWQGRALTNSSVY 60
QY 61 FVOYKIMFSCMSKSHQKPSGCHISCNPPGCRITLAKYQKQWKNKDCWGTQELSCDL 120
DB 61 FVOYKIMFSCMSKSHQKPSGCHISCNPPGCRITLAKYQKQWKNKDCWGTQELSCDL 120
QY 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
DB 121 TSSTDIQEPYGRVRAASAGSYSEWSMTPTPTWETKIDPPVNMITQVNGSLVILHA 180
QY 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
DB 181 PNLPRYQKKNVSIEDYELLYRVFIINNSLEKEQKVEGAHRAVEIHALTPHSSYCVV 240
QY 241 ASIYQMLDRRSORSERCEVEIP 263
DB 241 ASIYQMLDRRSORSERCEVEIP 263

```

RESULT 4

```

US-09-961-404-6
; Sequence 6, Application US/09961404
; Publication No. US2003022827A1
; GENERAL INFORMATION:
; APPLICANT: WEISS, BERTRAM
; APPLICANT: SABAT, ROBERT
; APPLICANT: ASADULLAH, KHUSRU
; APPLICANT: TOSCHI, LUISSELLA
; TITLE OF INVENTION: THREE NEW MEMBERS OF THE CYTOKINE RECEPTOR
; FILE REFERENCE: SCH-1788
; CURRENT APPLICATION NUMBER: US/09/961,404
; CURRENT FILING DATE: 2001-09-25
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: Patent in Ver. 2.1
; SEQ ID NO 6
; LENGTH: 263
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-961-404-6

```

```

Query Match 99.7%; Score 1428; DB 11; Length 263;
Best Local Similarity 99.6%; Pred. No. 1e-137;
Matches 262; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```